SEQUENCE LISTING

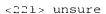
<110> Estell, David . Harding, Fiona <120> PROTEINS PRODUCING AN ALTERED IMMUNOGENIC RESPONSE AND METHODS OF MAKING AND USING THE SAME <130> A-68893/DJB/DAV <140> 09/500,135 <141> 2000-02-08 <150> 09/060,872 <151> 1998-04-15 <160> 236 <170> PatentIn Ver. 2.1 <210> 1 <211> 1495 <212> DNA <213> Bacillus amyloliquefaciens <220> <221> mat_peptide <222> (417)..(1495) <220> <221> CDS <222> (96)..(1244) <220> <221> unsure <222> (96)..(98) <223> The nnn at positions 96 through 98 represents gtg, which is to code for methionine. <220> <221> unsu**r**e <222> (582)..(584)

<220>

proline.

Xaa, which in a preferred embodiment (aat) is to code for asparagine, but which may also code for

<223> The nnn at positions 582 through 584 represents



- <222> (585)..(587)
- <223> The nnn at positions 585 through 587 represents
 Xaa, which in a preferred embodiment (cct) is to
 code for proline, but which may also code for
 asparagine.

<220>

- <221> unsure
- <222> (597)..(599)
- <223> The nnn at positions 597 to 599 represents Xaa, which in a preferred embodiment (aac) is to code for asparagine, but which may also code for aspartic acid.

<220>

- <221> unsure
- <222> (678)..(680)
- <223> The nnn at positions 678 through 680 represents Xaa, which in a preferred embodiment (gca) is to code for alanine, but which may also code for serine.

<220>

- <221> unsure
- <222> (681)..(683)
- <223> The nnn at positions 681 through 683 represents Xaa, which in a preferred embodiment (tca) is to code for serine, but which may also code for alanine.

<220>

- <221> unsure
- <222> (708)..(710)
- <223> The nnn at positions 708 through 710 represents Xaa, which in a preferred embodiment (gct) is to code for alanine, but which may also code for aspartic acid.

<220>

- <221> unsure
- <222> (711)..(713)
- <223> The nnn at positions 711 through 713 represents Xaa, which in a preferred embodiment (gac) is to code for aspartic acid, but which may also code for alanine.

<220>

<222>	unsure (888)(890) The nnn at positions 888 through 890 represents Xaa, which in a preferred embodiment (act) is to code for threonine, but which may also code for serine.														
<222>	<pre><220> <221> unsure <222> (891)(893) <223> The nnn at positions 891 through 893 represents</pre>														
<pre><220> <221> unsure <222> (1167)(1169) <223> The nnn at positions 1167 through 1169 represents</pre>															
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ttatt	ctgca	aatg	aaaa	aa aq	ggaga	agga	t aaa	_		Arg (aaa q Lys ^r	=	113
~ -	tc agt le Ser 00	_	_		_						-	_			161
	gc aca er Thr			-	-		_						_	_	209
	at att yr Ile	_				_		_	_	_	_	_	_	_	257

-50

aag aag aaa gat gtc att tct gaa aaa ggc ggg aaa gtg caa aag caa

Lys Lys Lys Asp Val Ile Ser Glu Lys Gly Gly Lys Val Gln Lys Gln

ttc aaa tat gta gac gca gct tca gct aca tta aac gaa aaa gct gta Phe Lys Tyr Val Asp Ala Ala Ser Ala Thr Leu Asn Glu Lys Ala Val

3

-45

305

-40





-35 -30 -25

Lys	Glu	Leu	Lys	Lys	Asp	Pro	Ser	Val	Ala	Tyr	Val	Glu	Glu	Asp	His	
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_	_				Ala	_										
- 5				-1	1				5	•	1			10		
	_		_	_	cac			-						_		497
Lys	Ala	Pro		Leu	His	Ser	Gln	_	Tyr	Thr	Gly	Ser		Val	Lys	
			15					20					25			
gta	gcg	gtt	atc	gac	agc	ggt	atc	gat	tct	tct	cat	cct	gat	tta	aag	545
Val	Ala	Val	Ile	Asp	Ser	Gly	Ile	Asp	Ser	Ser	His	Pro	Asp	Leu	Lys	
		30					35					40				
																503
_	_			_	agc	_	_			_						593
vai	45	GIY	GIÀ	Ala	Ser	мес 50	vaı	PIO	ser	Giu	55	лаа	лаа	Pne	GIII	
	43					50					55					
gac	nnn	aac	tct	cac	gga	act	cac	gtt	gcc	ggc	aca	gtt	gcg	gct	ctt	641
Asp	Xaa	Asn	Ser	His	Gly	Thr	His	Val	Ala	Gly	Thr	Val	Ala	Ala	Leu	
60					65					70					75	
aat	aac	tca	atc	aat	gta	tta	aac	att	aca	cca	aqc	nnn	nnn	ctt	tac	689
					Val			_			_					
				80			_		85					90		
_	_		_		ggt								_			737
Ala	Val	Lys		Leu	Gly	Xaa	Xaa	-	Ser	Gly	Gln	Tyr		Trp	Ile	
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Ile	Asn	Gly	Ile	Glu	Trp	Ala	Ile	Ala	Asn	Asn	Met	Asp	Val	Ile	Asn	
		110					115					120				
ata	200	ata	aac	aa a	cct	tat	aat	tat	act	act	t t a	222	aca	aca	att	833
					Pro											033
1100	125	Dea	Cry	Cry	110	130	OI y	DCI	1114	1114	135	270	1114	1114	• • • • • • • • • • • • • • • • • • • •	
gat	aaa	gcc	gtt	gca	tcc	ggc	gtc	gta	gtc	gtt	gcg	gca	gcc	ggt	aac	881
Asp	Lys	Ala	Val	Ala	Ser	Gly	Val	Val	Val	Val	Ala	Ala	Ala	Gly	Asn	
140					145					150					155	
gaa	aac	nnn	מממ	aac	agc	tca	age	aca	ata	aac	tac	cct	aat	aaa	tac	929
-	Gly				-		_									,.,





165 170 160 cet tet gte att gea gta gge get gtt gae age age aac caa aga gea Pro Ser Val Ile Ala Val Gly Ala Val Asp Ser Ser Asn Gln Arg Ala 175 180 tot tto toa ago gta gga cot gag ott gat gto atg goa cot ggo gta 1025 Ser Phe Ser Ser Val Gly Pro Glu Leu Asp Val Met Ala Pro Gly Val 190 195 tet ate caa age aeg ett eet gga aae aaa tae ggg geg tae aae ggt 1073 Ser Ile Gln Ser Thr Leu Pro Gly Asn Lys Tyr Gly Ala Tyr Asn Gly 205 215 210 acg tea atg gea tet eeg eac gtt gee gga geg get get ttg att ett 1121 Thr Ser Met Ala Ser Pro His Val Ala Gly Ala Ala Leu Ile Leu 220 225 230 235

tot aag cac cog aac tgg aca aac act caa gtc cgc agc agt tta nnn Ser Lys His Pro Asn Trp Thr Asn Thr Gln Val Arg Ser Ser Leu Xaa 245 240

aac acc act aca aaa ctt ggt gat tct ttc tac tat gga aaa ggg ctg Asn Thr Thr Lys Leu Gly Asp Ser Phe Tyr Tyr Gly Lys Gly Leu 255 260 265

atc aac gta cag gcg gca gct cag taa aacataaaaa accggccttg 1264 Ile Asn Val Gln Ala Ala Gln 270 275

geologicegg tittittatt titetteete egeatgitea ateegeteea taategaegg 1324 atggctccct ctgaaaattt taacgagaaa cggcgggttg acccggctca gtcccgtaac 1384 ggccaagtcc tgaaacgtct caatcgccgc ttcccggttt ccggtcagct caatgccgta 1444 acggtcggcg gcgttttcct gataccggga gacggcattc gtaatcggat c 1495

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<211> 382

<212> PRT

<213> Bacillus amyloliquefaciens

<400> 2

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Ser Th	r Met	Ser	Ala	Ala	Lys	Lys	Lys	Asp	Val	Ile	Ser	Glu	Lys	Gly
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Gly Ly	s Val	Gln	Lys	Gln	Phe	Lys	Tyr	Val	Asp	Ala	Ala	Ser	Ala	Thr
65				70					75					80
Leu As	n Glu	Lys		Val	Lys	Glu	Leu		Lys	Asp	Pro	Ser		Ala
			85					90					95	
Tyr Va	l Glu		Asp	His	Val	Ala		Ala	Tyr	Ala	Gln		Val	Pro
m 01		100	~ 1	- 1	_	- 1	105		-			110	~ 1	
Tyr Gl		ser	GIn	He	Lys		Pro	Ala	Leu	Hls		GIn	GIY	Tyr
mbs Cl	115	7.00	17al	T	17-1	120	₹ <i>₹</i> ~]	T10	7 00	Cox	125	T 10	7 ~~	Cor
Thr Gl		ASII	vaı	пуъ	135	Ala	vaı	TIE	Asp	140	GIY	116	нър	ser
Ser Hi		Δsn	T.e.u	Lve		Δla	Glv	Glv	Δla		Met	Val	Pro	Ser
145	5 110	пър	пси	150	vai	HIU	Gry	Gry	155	DCI	1100	Val	110	160
Glu Th	r Xaa	Xaa	Phe		Asp	Xaa	Asn	Ser		Glv	Thr	His	Val	
			165					170		1			175	
Gly Th	r Val	Ala	Ala	Leu	Asn	Asn	Ser	Ile	Gly	Val	Leu	Gly	Val	Ala
		180					185					190		
Pro Se	r Xaa	Xaa	Leu	Tyr	Ala	Val	Lys	Val	Leu	Gly	Xaa	Xaa	Gly	Ser
	195					200					205			
Gly Gl	n Tyr	Ser	Trp	Ile	Ile	Asn	Gly	Ile	Glu	Trp	Ala	Ile	Ala	Asn
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Asn Me	t Asp	Val	Ile	Asn	Met	Ser	Leu	Gly	Gly	Pro	Ser	Gly	Ser	Ala
225				230					235					240
Ala Le	u Lys	Ala		Val	Asp	Lys	Ala		Ala	Ser	Gly	Val		Val
** 7 7 7 7		2.7	245	_	a.	~ 3	77	250	~ 1			a	255	** 3
Val Al	а Ата	260	GIY	ASI	GIU	GIY	265	хаа	GIY	ser	ser	270	Thr	vaı
Gly Ty	r Pro		Lve	Tur	Pro	Ser		Tla	בומ	Val	Glv		Val	Aen
Cry ry	275	Ory	Lys	1 y 1	110	280	val	110	7114	vai	285	7114	Vai	пър
Ser Se		Gln	Arq	Ala	Ser		Ser	Ser	Val	Glv		Glu	Leu	Asp
29			5		295					300				_
Val Me	t Ala	Pro	Gly	Val	Ser	Ile	Gln	Ser	Thr	Leu	Pro	Gly	Asn	Lys
305				310					315					320
Tyr Gl	y Ala	Tyr	Asn	Gly	Thr	Ser	Met	Ala	Ser	Pro	His	Val	Ala	Gly
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Ala Al	a Ala	Leu	Ile	Leu	Ser	Lys	His	Pro	Asn	Trp	Thr	Asn	Thr	Gln
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Val Ar	g Ser	Ser	Leu	Xaa	Asn	Thr	Thr	Thr	Lys	Leu	Gly	Asp	Ser	Phe
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<213> Bacillus amyloliquefaciens

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Ser Gly Ile Asp Ser Ser His Pro Asp Leu Lys Val Ala Gly Gly Ala 35 40 45

Ser Met Val Pro Ser Glu Thr Asn Pro Phe Gln Asp Asn Asn Ser His
50 55 60

Gly Thr His Val Ala Gly Thr Val Ala Ala Leu Asn Asn Ser Ile Gly 65 70 75 80

Val Leu Gly Val Ala Pro Ser Ala Ser Leu Tyr Ala Val Lys Val Leu 85 90 95

Gly Ala Asp Gly Ser Gly Gln Tyr Ser Trp Ile Ile Asn Gly Ile Glu 100 105 110

Trp Ala Ile Ala Asn Asn Met Asp Val Ile Asn Met Ser Leu Gly Gly
115 120 125

Pro Ser Gly Ser Ala Ala Leu Lys Ala Ala Val Asp Lys Ala Val Ala 130 135 140

Ser Gly Val Val Val Ala Ala Ala Gly Asn Glu Gly Thr Ser Gly
145 150 155 160

Ser Ser Ser Thr Val Gly Tyr Pro Gly Lys Tyr Pro Ser Val Ile Ala 165 170 175

Val Gly Ala Val Asp Ser Ser Asn Gln Arg Ala Ser Phe Ser Ser Val 180 185 190

Gly Pro Glu Leu Asp Val Met Ala Pro Gly Val Ser Ile Gln Ser Thr 195 200 205

Leu Pro Gly Asn Lys Tyr Gly Ala Tyr Asn Gly Thr Ser Met Ala Ser 210 215 220



Pro His Val Ala Gly Ala Ala Ala Leu Ile Leu Ser Lys His Pro Asn 225 230 235 240

Trp Thr Asn Thr Gln Val Arg Ser Ser Leu Glu Asn Thr Thr Lys
245 250 255

Leu Gly Asp Ser Phe Tyr Tyr Gly Lys Gly Leu Ile Asn Val Gln Ala 260 265 270

Ala Ala Gln 275

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<211> 275

<212> PRT

<213> Bacillus subtilis

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His Ser Gln Gly Tyr Thr Gly Ser Asn Val Lys Val Ala Val Ile Asp 20 25 30

Ser Gly Ile Asp Ser Ser His Pro Asp Leu Asn Val Arg Gly Gly Ala 35 40 45

Ser Phe Val Pro Ser Glu Thr Asn Pro Tyr Gln Asp Gly Ser Ser His 50 55 60

Gly Thr His Val Ala Gly Thr Ile Ala Ala Leu Asn Asn Ser Ile Gly 65 70 75 80

Val Leu Gly Val Ser Pro Ser Ala Ser Leu Tyr Ala Val Lys Val Leu 85 90 95

Asp Ser Thr Gly Ser Gly Gln Tyr Ser Trp Ile Ile Asn Gly Ile Glu 100 105 110

Trp Ala Ile Ser Asn Asn Met Asp Val Ile Asn Met Ser Leu Gly Gly
115 120 125

Pro Thr Gly Ser Thr Ala Leu Lys Thr Val Val Asp Lys Ala Val Ser 130 135 140

Ser Gly Ile Val Val Ala Ala Ala Gly Asn Glu Gly Ser Ser Gly
145 150 155 160



Ser Thr Ser Thr Val Gly Tyr Pro Ala Lys Tyr Pro Ser Thr Ile Ala 165 170 175

Val Gly Ala Val Asn Ser Ser Asn Gln Arg Ala Ser Phe Ser Ser Ala 180 185 190

Gly Ser Glu Leu Asp Val Met Ala Pro Gly Val Ser Ile Gln Ser Thr 195 200 205

Leu Pro Gly Gly Thr Tyr Gly Ala Tyr Asn Gly Thr Ser Met Ala Thr 210 215 220

Pro His Val Ala Gly Ala Ala Ala Leu Ile Leu Ser Lys His Pro Thr 225 230 235 240

Trp Thr Asn Ala Gln Val Arg Asp Arg Leu Glu Ser Thr Ala Thr Tyr 245 250 255

Leu Gly Asn Ser Phe Tyr Tyr Gly Lys Gly Leu Ile Asn Val Gln Ala 260 265 270

Ala Ala Gln 275

<210> 5

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<213> Bacillus licheniformis

<400> 5

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Gln Ala Gln Gly Phe Lys Gly Ala Asn Val Lys Val Ala Val Leu Asp 20 25 30

Thr Gly Ile Gln Ala Ser His Pro Asp Leu Asn Val Val Gly Gly Ala 35 40 45

Ser Phe Val Ala Gly Glu Ala Tyr Asn Thr Asp Gly Asn Gly His Gly 50 55 60

Thr His Val Ala Gly Thr Val Ala Ala Leu Asp Asn Thr Thr Gly Val 65 70 75 80

Leu Gly Val Ala Pro Ser Val Ser Leu Tyr Ala Val Lys Val Leu Asn

85 90 95

Ser Ser Gly Ser Gly Ser Tyr Ser Gly Ile Val Ser Gly Ile Glu Trp 100 105 110

Ala Thr Thr Asn Gly Met Asp Val Ile Asn Met Ser Leu Gly Gly Ala 115 120 125

Ser Gly Ser Thr Ala Met Lys Gln Ala Val Asp Asn Ala Tyr Ala Arg 130 135 140

Gly Val Val Val Val Ala Ala Ala Gly Asn Ser Gly Asn Ser Gly Ser 145 150 155 160

Thr Asn Thr Ile Gly Tyr Pro Ala Lys Tyr Asp Ser Val Ile Ala Val 165 170 175

Gly Ala Val Asp Ser Asn Ser Asn Arg Ala Ser Phe Ser Ser Val Gly
180 185 190

Ala Glu Leu Glu Val Met Ala Pro Gly Ala Gly Val Tyr Ser Thr Tyr 195 200 205

Pro Thr Asn Thr Tyr Ala Thr Leu Asn Gly Thr Ser Met Ala Ser Pro 210 215 220

His Val Ala Gly Ala Ala Ala Leu Ile Leu Ser Lys His Pro Asn Leu 225 230 235 240

Ser Ala Ser Gln Val Arg Asn Arg Leu Ser Ser Thr Ala Thr Tyr Leu 245 250 255

Gly Ser Ser Phe Tyr Tyr Gly Lys Gly Leu Ile Asn Val Glu Ala Ala 260 265 270

Ala Gln

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<213> Bacillus lentus

<400> 6

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35 40 45

Phe Val Pro Gly Glu Pro Ser Thr Gln Asp Gly Asn Gly His Gly Thr 50 55 60

His Val Ala Gly Thr Ile Ala Ala Leu Asn Asn Ser Ile Gly Val Leu 65 70 75 80

Gly Val Ala Pro Ser Ala Glu Leu Tyr Ala Val Lys Val Leu Gly Ala 85 90 95

Ser Gly Ser Gly Ser Val Ser Ser Ile Ala Gln Gly Leu Glu Trp Ala 100 105 110

Gly Asn Asn Gly Met His Val Ala Asn Leu Ser Leu Gly Ser Pro Ser 115 120 125

Pro Ser Ala Thr Leu Glu Gln Ala Val Asn Ser Ala Thr Ser Arg Gly 130 135 140

Val Leu Val Val Ala Ala Ser Gly Asn Ser Gly Ala Gly Ser Ile Ser 145 150 155 160

Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val Gly Ala Thr Asp Gln
165 170 175

Asn Asn Asn Arg Ala Ser Phe Ser Gln Tyr Gly Ala Gly Leu Asp Ile 180 185 190

Val Ala Pro Gly Val Asn Val Gln Ser Thr Tyr Pro Gly Ser Thr Tyr
195 200 205

Ala Ser Leu Asn Gly Thr Ser Met Ala Thr Pro His Val Ala Gly Ala 210 215 220

Ala Ala Leu Val Lys Gln Lys Asn Pro Ser Trp Ser Asn Val Gln Ile 225 230 235 240

Arg Asn His Leu Lys Asn Thr Ala Thr Ser Leu Gly Ser Thr Asn Leu 245 250 255

Tyr Gly Ser Gly Leu Val Asn Ala Glu Ala Ala Thr Arg \$260\$

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<210> 83

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<210> 96 <211> 15 <212> PRT <213> Artificial Sequence

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Arg Ala Ser Leu Ser Leu Gly Ser Gly Phe Trp His Ala Thr Gly
1 5 10 15

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1 5 10 15

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Arg His Ser Ser Arg Arg Leu Leu Arg Ala Ile Pro Arg Gln Val

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                                     10
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                  5
                                    10
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Ala Ile Pro Arg Gln Val Ala Gln Thr Leu Gln Ala Asp Val Leu
                 5
                                     10
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<211> 15
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<400> 106

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1 5 10 15

<210> 107

<211> 15

<212> PRT

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<223> Description of Artificial Sequence: Synthetic

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1 5 10 15

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<223> Description of Artificial Sequence: Synthetic

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1 5 10 15

<210> 109

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<400> 109

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          5
                                     10
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                                                         15
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Lys His Pro His Phe Lys Asn Val Lys Glu Arg Thr Asn Trp Thr 1 5 10 15

<210> 118

<211> 15

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<223> Description of Artificial Sequence: Synthetic

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His Phe Lys Asn Val Lys Glu Arg Thr Asn Trp Thr Asn Glu Arg
1 5 10 15

<210> 119

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Asn Val Lys Glu Arg Thr Asn Trp Thr Asn Glu Arg Thr Leu Asp
1 5 10 15

<210> 120

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<223> Description of Artificial Sequence: Synthetic

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Glu Arg Thr Asn Trp Thr Asn Glu Arg Thr Leu Asp Asp Gly Leu

1 5 10 15

<210> 121

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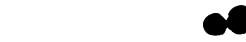
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<223> Description of Artificial Sequence: Synthetic
<400> 128
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Ile Ala Ser Met Arg Glu Cys Gln Gly Phe Ala Pro Asp Ala Glu



1 5 10 15

<210> 129

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<400> 129

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1 5 10 15

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<210> 131

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<223> Description of Artificial Sequence: Synthetic

<400> 131

Phe Ala Pro Asp Ala Glu Leu His Ile Phe Arg Val Phe Thr Asn 1 5 10 15

<210> 132

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                  5
                                     10
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                  5
                                     10
                                                          15
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<212> PRT
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Ser Trp Phe Leu Asp Ala Phe Asn Tyr Ala Ile Leu Lys Lys Ile
                 5
                                     10
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                                     10
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<400> 148
Pro Phe Val Asp Lys Val Trp Glu Leu Thr Ala Asn Asn Val Ile
                 5
                                     10
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Asp Lys Val Trp Glu Leu Thr Ala Asn Asn Val Ile Met Val Ser
                  5
                                     10
 1
                                                          15
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Trp Glu Leu Thr Ala Asn Asn Val Ile Met Val Ser Ala Ile Gly
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                                    10
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                                      10
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<400> 152
Asn Val Ile Met Val Ser Ala Ile Gly Asn Asp Gly Pro Leu Tyr
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<210> 154
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Met Asp Val Ile Gly Val Gly Gly Ile Asp Phe Glu Asp Asn Ile
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Ala Arg Phe Ser Ser Arg Gly Met Thr Thr Trp Glu Leu Pro Gly
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<210> 166

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<400> 166

Ser Ser Arg Gly Met Thr Thr Trp Glu Leu Pro Gly Gly Tyr Gly
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<210> 167

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Arg Met Lys Pro Asp Ile Val Thr Tyr Gly Ala Gly Val Arg Gly
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Pro Asp Ile Val Thr Tyr Gly Ala Gly Val Arg Gly Ser Gly Val
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1 5 10 15

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<210> 178

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<210> 179

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<210> 186
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Arg Glu Leu Val Asn Pro Ala Ser Met Lys Gln Ala Leu Ile Ala
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Asp Leu Leu Arg Ala Tyr Gln Ile Leu Asn Ser Tyr Lys Pro Gln
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Gln Ile Leu Asn Ser Tyr Lys Pro Gln Ala Ser Leu Ser Pro Ser
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Asn Ser Tyr Lys Pro Gln Ala Ser Leu Ser Pro Ser Tyr Ile Asp
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<210> 202
<211> 15
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Lys Pro Gln Ala Ser Leu Ser Pro Ser Tyr Ile Asp Leu Thr Glu
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<211> 15
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<400> 203
Ala Ser Leu Ser Pro Ser Tyr Ile Asp Leu Thr Glu Cys Pro Tyr
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<210> 204
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<212> PRT
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Ser Pro Ser Tyr Ile Asp Leu Thr Glu Cys Pro Tyr Met Trp Pro
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Tyr Ile Asp Leu Thr Glu Cys Pro Tyr Met Trp Pro Tyr Cys Ser
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Glu Val Asp Asn Trp Arg Ile Ile Pro Arg Asn Asn Pro Ser Ser Asp

Tyr Pro Ser Asp Phe Glu Val Ile Gln Ile Lys Glu Lys Gln Lys Ala

105

90

110

85

Gly	Leu	Leu 115	Thr	Leu	Glu	Asp	His 120	Pro	Asn	Ile	Lys	Arg 125	Val	Thr	Pro
Gln	Arg 130	Lys	Val	Phe	Arg	Ser 135	Leu	Lys	Tyr	Ala	Glu 140	Ser	Asp	Pro	Thr
Val 145	Pro	Cys	Asn	Glu	Thr 150	Arg	Trp	Ser	Gln	Lys 155	Trp	Gln	Ser	Ser	Arg 160
Pro	Leu	Arg	Arg	Ala 165	Ser	Leu	Ser	Leu	Gly 170	Ser	Gly	Phe	Trp	His 175	Ala
Thr	Gly	Arg	His 180	Ser	Ser	Arg	Arg	Leu 185	Leu	Arg	Ala	Ile	Pro 190	Arg	Gln
Val	Ala	Gln 195	Thr	Leu	Gln	Ala	Asp 200	Val	Leu	Trp	Gln	Met 205	Gly	Tyr	Thr
Gly	Ala 210	Asn	Val	Arg	Val	Ala 215	Val	Phe	Asp	Thr	Gly 220	Leu	Ser	Glu	Lys
His 225	Pro	His	Phe	Lys	Asn 230	Val	Lys	Glu	Arg	Thr 235	Asn	Trp	Thr	Asn	Glu 240
Arg	Thr	Leu	Asp	Asp 245	Gly	Leu	Gly	His	Gly 250	Thr	Phe	Val	Ala	Gly 255	Val
Ile	Ala	Ser	Met 260	Arg	Glu	Cys	Gln	Gly 265	Phe	Ala	Pro	Asp	Ala 270	Glu	Leu
His	Ile	Phe 275	Arg	Val	Phe	Thr	Asn 280	Asn	Gln	Val	Ser	Tyr 285	Thr	Ser	Trp
Phe	Leu 290	Asp	Ala	Phe	Asn	Tyr 295	Ala	Ile	Leu	Lys	Lys 300	Ile	Asp	Val	Leu
Asn 305	Leu	Ser	Ile	Gly	Gly 310	Pro	Asp	Phe	Met	Asp 315	His	Pro	Phe	Val	Asp 320
Lys	Val	Trp	Glu	Leu 325	Thr	Ala	Asn	Asn	Val 330	Ile	Met	Val	Ser	Ala 335	Ile
Gly	Asn	Asp	Gly 340	Pro	Leu	Tyr	Gly	Thr 345	Leu	Asn	Asn	Pro	Ala 350	Asp	Gln
Met	Asp	Val 355	Ile	Gly	Val	Gly	Gly 360	Ile	Asp	Phe	Glu	Asp 365	Asn	Ile	Ala

Arg Phe Ser	Ser Arg	Gly Met		nr Trp G	lu Leu 380	Pro	Gly	Gly	Tyr
Gly Arg Met	Lys Pro	Asp Ile	Val Th	-	ly Ala 95	Gly	Val	Arg	Gly 400
Ser Gly Val	Lys Gly 405	Gly Cys	Arg Al	la Leu S 410	er Gly	Thr	Ser	Val 415	Ala
Ser Pro Val	Val Ala 420	Gly Ala	Val Th		eu Val	Ser	Thr 430	Val	Gln
Lys Arg Glu		Asn Pro	Ala Se	er Met L	ys Gln	Ala 445	Leu	Ile	Ala
Ser Ala Arç 450	Arg Leu	Pro Gly 455		sn Met P	he Glu 460	Gln	Gly	His	Gly
Lys Leu Asp 465	Leu Leu	Arg Ala	Tyr Gl		eu Asn 75	Ser	Tyr	Lys	Pro 480
Gln Ala Sen	Leu Ser 485	Pro Ser	Tyr Il	le Asp L 490	eu Thr	Glu	Cys	Pro 495	Tyr
Met Trp Pro	Tyr Cys 500	Ser Gln	Pro Il	-	yr Gly	_	Met 510	Pro	Thr
Val Val Ası		Ile Leu	Asn Gl	ly Met G	ly Val	Thr 525	Gly	Arg	Ile
Val Asp Lys	Pro Asp	Trp Gln 535	-	yr Leu P	ro Gln 540	Asn	Gly	Asp	Asn
Ile Glu Val	Ala Phe	Ser Tyr 550	Ser Se		eu Trp 55	Pro	Trp	Ser	Gly 560
Tyr Leu Ala	Ile Ser 565	Ile Ser	Val Th	ir Lys Ly 570	ys Ala	Ala		Trp 575	Glu
Gly Ile Ala	Gln Gly	His Val	Met Il		al Ala		Pro 590	Ala	Glu
Thr Glu Sen	•	Gly Ala	Glu Gl 600	n Thr Se	er Thr	Val 605	Lys	Leu	Pro

Trp 625	Asp	Gln	Tyr	His	Asn 630	Leu	Arg	Tyr	Pro	Pro 635	Gly	Tyr	Phe	Pro	Arg 640
Asp	Asn	Leu	Arg	Met 645	Lys	Asn	Asp	Pro	Leu 650	Asp	Trp	Asn	Gly	Asp 655	His
Ile	His	Thr	Asn 660	Phe	Arg	Asp	Met	Tyr 665	Gln	His	Leu	Arg	Ser 670	Met	Gly
Tyr	Phe	Val 675	Glu	Val	Leu	Gly	Ala 680	Pro	Phe	Thr	Cys	Phe 685	Asp	Ala	Ser
Gln	Tyr 690	Gly	Thr	Leu	Leu	Met 695	Val	Asp	Ser	Glu	Glu 700	Glu	Tyr	Phe	Pro
Glu 705	Glu	Ile	Ala	Lys	Leu 710	Arg	Arg	Asp	Val	Asp 715	Asn	Gly	Leu	Ser	Leu 720
Val	Ile	Phe	Ser	Asp 725	Trp	Tyr	Asn	Thr	Ser 730	Val	Met	Arg	Lys	Val 735	Lys
Phe	Tyr	Asp	Glu 740	Asn	Thr	Arg	Gln	Trp 745	Trp	Met	Pro	Asp	Thr 750	Gly	Gly
Ala	Asn	Ile 755	Pro	Ala	Leu	Asn	Glu 760	Leu	Leu	Ser	Val	Trp 765	Asn	Met	Gly
Phe	Ser 770	Asp	Gly	Leu	Tyr	Glu 775	Gly	Glu	Phe	Thr	Leu 780	Ala	Asn	His	Asp
Met 785	Tyr	Tyr	71-	_											
			AIA	Ser	Gly 790	Cys	Ser	Ile	Ala	Lys 795	Phe	Pro	Glu	Asp	Gly 800
Val	Val	Ile			790	-				- 795				Asp Leu 815	800
			Thr	Gln 805	790 Thr	Phe	Lys	Asp	Gln 810	795 Gly	Leu	Glu	Val	Leu	800 Lys
Gln	Glu	Thr	Thr Ala 820	Gln 805 Val	790 Thr	Phe Glu	Lys Asn	Asp Val 825	Gln 810 Pro	795 Gly	Leu Leu	Glu Gly	Val Leu 830	Leu 815	800 Lys Gln
Gln	Glu	Thr Ala 835	Thr Ala 820 Glu	Gln 805 Val	790 Thr Val	Phe Glu Gly	Lys Asn Arg 840	Asp Val 825 Ile	Gln 810 Pro Val	795 Gly Ile	Leu Leu Tyr	Glu Gly Gly 845	Val Leu 830 Asp	Leu 815 Tyr	800 Lys Gln Asn

His Ser Gly Asn Arg Gln Arg Pro Pro Ser Gly Ala Gly Ser Val Thr 885 890 Pro Glu Arg Met Glu Gly Asn His Leu His Arg Tyr Ser Lys Val Leu 900 905 Glu Ala His Leu Gly Asp Pro Lys Pro Arg Pro Leu Pro Ala Cys Pro 920 Arg Leu Ser Trp Ala Lys Pro Gln Pro Leu Asn Glu Thr Ala Pro Ser 930 935 940 Asn Leu Trp Lys His Gln Lys Leu Leu Ser Ile Asp Leu Asp Lys Val 950 955 Val Leu Pro Asn Phe Arg Ser Asn Arg Pro Gln Val Arg Pro Leu Ser 965 970 Pro Gly Glu Ser Gly Ala Trp Asp Ile Pro Gly Gly Ile Met Pro Gly 980 985 Arg Tyr Asn Gln Glu Val Gly Gln Thr Ile Pro Val Phe Ala Phe Leu 995 1000 1005 Gly Ala Met Val Val Leu Ala Phe Phe Val Val Gln Ile Asn Lys Ala 1010 1015 1020 Lys Ser Arg Pro Lys Arg Arg Lys Pro Arg Val Lys Arg Pro Gln Leu 1025 1035 1040

Met Gln Gln Val His Pro Pro Lys Thr Pro Ser Val

1045 1050

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Trp Gln Met Gly Tyr Thr Gly Ala Asn Val Arg Val Ala Val Phe Asp
20 25 30

Thr Gly Leu Ser Glu Lys His Pro His Phe Lys Asn Val Lys Glu Arg

Thr	Asn 50	Trp	Thr	Asn	Glu	Arg 55	Thr	Leu	Asp	Asp	Gly 60	Leu	Gly	His	Gly
Thr 65	Phe	Val	Ala	Gly	Val 70	Ile	Ala	Ser	Met	Arg 75	Glu	Cys	Gln	Gly	Phe 80
Ala	Pro	Asp	Ala	Glu 85	Leu	His	Ile	Phe	Arg 90	Val	Phe	Thr	Asn	Asn 95	Gln
Val	Ser	Tyr	Thr 100	Ser	Trp	Phe	Leu	Asp 105	Ala	Phe	Asn	Tyr	Ala 110	Ile	Leu
Lys	Lys	Ile 115	Asp	Val	Leu	Asn	Leu 120	Ser	Ile	Gly	Gly	Pro 125	Asp	Phe	Met
Asp	His 130	Pro	Phe	Val	Asp	Lys 135	Val	Trp	Glu	Leu	Thr 140	Ala	Asn	Asn	Val
Ile 145	Met	Val	Ser	Ala	Ile 150	Gly	Asn	Asp	Gly	Pro 155	Leu	Tyr	Gly	Thr	Leu 160
Asn	Asn	Pro	Ala	Asp 165	Gln	Met	Asp	Val	Ile 170	Gly	Val	Gly	Gly	Ile 175	Asp
Phe	Glu	Asp	Asn 180	Ile	Ala	Arg	Phe	Ser 185	Ser	Arg	Gly	Met	Thr 190	Thr	Trp
Glu	Leu	Pro 195	Gly	Gly	Tyr	Gly	Arg 200	Met	Lys	Pro	Asp	Ile 205	Val	Thr	Tyr
Gly	Ala 210	Gly	Val	Arg	Gly	Ser 215	Gly	Val	Lys	Gly	Gly 220	Cys	Arg	Ala	Leu
Ser 225	Gly	Thr	Ser	Val	Ala 230	Ser	Pro	Val	Val	Ala 235	Gly	Ala	Val	Thr	Leu 240
Leu	Val	Ser	Thr	Val 245	Gln	Lys	Arg	Glu	Leu 250	Val	Asn	Pro	Ala	Ser 255	Met
Lys	Gln	Ala	Leu 260	Ile	Ala	Ser	Ala	Arg 265	Arg	Leu	Pro	Gly	Val 270	Asn	Met
Phe	Glu	Gln	Gly	His	Gly	Lys	Leu								

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<400> 211
Ala Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val
                 5
                                     10
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Gly Ala Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val
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                 5
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Gly Ser Ile Ser Tyr Pro Ala Arg Ala Ala Asn Ala Met Ala Val
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Gly Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Ala Ala Met Ala Val
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Val Phe Ser Cys Asn Ala Asn Phe Gln Arg Ile Thr Asp Phe Asp Ala

50 55 60

Lys Ser Gly Cys Glu Pro Gly Gly Val Ala Tyr Ser Cys Ala Asp Gln 65 70 75 80

Thr Pro Trp Ala Val Asn Asp Asp Phe Ala Leu Gly Phe Ala Ala Thr 85 90 95

Ser Ile Ala Gly Ser Asn Glu Ala Gly Trp Cys Cys Ala Cys Tyr Glu 100 105 110

Leu Thr Phe Thr Ser Gly Pro Val Ala Gly Lys Lys Met Val Val Gln
115 120 125

Ser Thr Ser Thr Gly Gly Asp Leu Gly Ser Asn His Phe Asp Leu Asn 130 135 140

Gly Gly Leu Pro Gly Gln Arg Tyr Gly Gly Ile Ser Ser Arg Asn Glu 165 170 175

Cys Asp Arg Phe Pro Asp Ala Leu Lys Pro Gly Cys Tyr Trp Arg Phe 180 185 190

Asp Trp Phe Lys Asn Ala Asp Asn Pro Ser Phe Ser Phe Arg Gln Val
195 200 205

Gln Cys Pro Ala Glu Leu Val Ala Arg Thr Gly Cys Arg Arg Asn Asp 210 215 220

Asp Gly Asn Phe Pro Ala Val Gln Ile Pro Ser Ser Ser Thr Ser Ser 225 230 235 240

Pro Val Asn Gln Pro Thr Ser Thr Ser Thr Thr Ser Thr Ser Thr Thr 245 250 255

Ser Ser Pro Pro Val Gln Pro Thr Thr Pro Ser Gly Cys Thr Ala Glu 260 265 270

Arg Trp Ala Gln 275

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<211> 18

<212> PRT

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Gly Asn Leu Asn Phe Asp Leu Lys Glu Ile Asn Asp Ile Cys Ser Gly

115 120 125

Cys Arg Gly His Asp Gly Phe Thr Ser Ser Trp Arg Ser Val Ala Asp 130 135 140

Arg Val Val Phe Thr Gly His Ser Leu Gly Gly Ala Leu Ala Thr Val 165 170 175

Ala Gly Ala Asp Leu Arg Gly Asn Gly Tyr Asp Ile Asp Val Phe Ser 180 185 190

Tyr Gly Ala Pro Arg Val Gly Asn Arg Ala Phe Ala Glu Phe Leu Thr
195 200 205

Val Gln Thr Gly Gly Thr Leu Tyr Arg Ile Thr His Thr Asn Asp Ile 210 215 220

Val Pro Arg Leu Pro Pro Arg Glu Phe Gly Tyr Ser His Ser Ser Pro 225 230 235 240

Glu Tyr Trp Ile Lys Ser Gly Thr Leu Val Pro Val Thr Arg Asn Asp 245 250 255

Ile Val Lys Ile Glu Gly Ile Asp Ala Thr Gly Gly Asn Asn Gln Pro
260 265 270

Asn Ile Pro Asp Ile Pro Ala His Leu Trp Tyr Phe Gly Leu Ile Gly 275 280 285

Thr Cys Leu 290

<210> 228

<211> 15

<212> PRT

<213> Streptomyces plicatus

<400> 228

Ile Lys Val Leu Leu Ser Val Leu Gly Asn His Gln Gly Ala Gly
1 5 10 15

<210> 229

<211> 313

-71	1 2	~ '	ÞΕ	ידי כ
< 1	- 2	>	r	< 1

<213> Streptomyces plicatus

<400> 229

- Met Phe Thr Pro Val Arg Arg Val Arg Thr Ala Ala Leu Ala Leu
 1 5 10 15
- Ser Ala Ala Ala Leu Val Leu Gly Ser Thr Ala Ala Ser Gly Ala 20 25 30
- Ser Ala Thr Pro Ser Pro Ala Pro Ala Pro Ala Pro Ala Pro Val Lys 35 40 45
- Gln Gly Pro Thr Ser Val Ala Tyr Val Glu Val Asn Asn Asn Ser Met 50 55 60
- Leu Asn Val Gly Lys Tyr Thr Leu Ala Asp Gly Gly Asn Ala Phe
 65 70 75 80
- Asp Val Ala Val Ile Phe Ala Ala Asn Ile Asn Tyr Asp Thr Gly Thr
 85 90 95
- Lys Thr Ala Tyr Leu His Phe Asn Glu Asn Val Gln Arg Val Leu Asp 100 105 110
- Asn Ala Val Thr Gln Ile Arg Pro Leu Gln Gln Gln Gly Ile Lys Val 115 120 125
- Leu Leu Ser Val Leu Gly Asn His Gln Gly Ala Gly Phe Ala Asn Phe 130 135 140
- Val Ala Lys Tyr Gly Leu Asp Gly Val Asp Phe Asp Asp Glu Tyr Ala 165 170 175
- Glu Tyr Gly Asn Asn Gly Thr Ala Gln Pro Asn Asp Ser Ser Phe Val 180 185 190
- His Leu Val Thr Ala Leu Arg Ala Asn Met Pro Asp Lys Ile Ile Ser 195 200 205
- Leu Tyr Asn Ile Gly Pro Ala Ala Ser Arg Leu Ser Tyr Gly Gly Val 210 215 220
- Asp Val Ser Asp Lys Phe Asp Tyr Ala Trp Asn Pro Tyr Tyr Gly Thr 225 230 235 240

Trp Gln Val Pro Gly Ile Ala Leu Pro Lys Ala Gln Leu Ser Pro Ala 245 250 255

Ala Val Glu Ile Gly Arg Thr Ser Arg Ser Thr Val Ala Asp Leu Ala 260 265 270

Arg Arg Thr Val Asp Glu Gly Tyr Gly Val Tyr Leu Thr Tyr Asn Leu 275 280 285

Asp Gly Gly Asp Arg Thr Ala Asp Val Ser Ala Phe Thr Arg Glu Leu 290 295 300

Tyr Gly Ser Glu Ala Val Arg Thr Pro 305 310

<210> 230

<211> 15

<212> PRT

<213> Bacillus amyloliquefaciens

<400> 230

Gly Thr Val Ala Ala Leu Asn Asn Ser Ile Gly Val Leu Gly Val
1 5 10 15

<210> 231

<211> 15

<212> PRT

<213> Bacillus amyloliquefaciens

<400> 231

Asn Gly Ile Glu Trp Ala Ile Ala Asn Asn Met Asp Val Ile Asn 1 5 10 15

<210> 232

<211> 15

<212> PRT

<213> Bacillus lentus

<400> 232

Thr Gly Ser Gly Val Lys Val Ala Val Leu Asp Thr Gly Ile Ser
1 5 10 15

<210> 233

<211> 15 <212> PRT <213> Bacillus lentus <400> 233 Ser Ala Glu Leu Tyr Ala Val Lys Val Leu Gly Ala Ser Gly Ser 5 10 <210> 234 <211> 17 <212> PRT <213> Bacillus lentus <400> 234 Gly Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val Gly 10 5 Ala <210> 235 <211> 15 <212> PRT <213> Bacillus lentus <400> 235 Gly Ala Gly Leu Asp Ile Val Ala Pro Gly Val Asn Val Gln Ser 5 10 15 <210> 236 <211> 272 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Hybrid of Bacillus lentus and Bacillus amyloliquefaciens Ala Gln Ser Val Pro Trp Gly Ile Ser Arg Val Gln Ala Pro Ala Ala 5 10

20

His Asn Arg Gly Leu Thr Gly Ser Gly Val Lys Val Ala Val Leu Asp

25

Thr	GIY	35	Ser	Thr	His	Pro	Asp 40	Leu	Asn	11e	Arg	45	GIY	АІА	ser
Phe	Val 50	Pro	Gly	Glu	Pro	Ser 55	Thr	Gln	Asp	Gly	Asn 60	Gly	His	Gly	Thr
His 65	Val	Ala	Gly	Thr	Ile 70	Ala	Ala	Leu	Asn	Asn 75	Ser	Ile	Gly	Val	Leu 80
Gly	Val	Ala	Pro	Ser 85	Ala	Glu	Leu	Tyr	Ala 90	Val	Lys	Val	Leu	Gly 95	Ala
Ser	Gly	Ser	Gly 100	Ser	Val	Ser	Ser	Ile 105	Ala	Gln	Gly	Leu	Glu 110	Trp	Ala
Gly	Asn	Asn 115	Gly	Met	His	Val	Ile 120	Asn	Met	Ser	Leu	Gly 125	Gly	Ser	Gly
Ser	Ala 130	Ala	Leu	Lys	Ala	Ala 135	Val	Asp	Lys	Ala	Val 140	Ala	Ser	Gly	Val
Val 145	Val	Val	Ala	Ala	Ala 150	Gly	Asn	Glu	Gly	Thr 155	Ser	Gly	Ser	Ser	Ser 160
Thr	Val	Gly	Tyr	Pro 165	Gly	Lys	Tyr	Pro	Ser 170	Val	Ile	Ala	Val	Gly 175	Ala
Val	Asp	Ser	Ser 180	Asn	Gln	Arg	Ala	Ser 185	Phe	Ser	Ser	Val	Gly 190	Pro	Glu
Leu	Asp	Val 195	Met	Ala	Pro	Gly	Val 200	Ser	Ile	Gln	Ser	Thr 205	Leu	Pro	Gly
Asn	Lys 210	Tyr	Gly	Ala	Tyr	Asn 215	Gly	Thr	Ser	Met	Ala 220	Ser	Pro	His	Val
225	-				230	Ile			-	235			_		240
Thr	Gln	Val	Arg	Ser 245	Ser	Leu	Glu	Asn	Thr 250	Thr	Thr	Lys	Leu	Gly 255	Asp

Thr Gly Ile Ser Thr His Pro Asp Leu Asn Ile Arg Gly Gly Ala Ser

Ser Phe Tyr Tyr Gly Lys Gly Leu Ile Asn Val Gln Ala Ala Gln

270